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Timestamp: [year=2007; month=12; day=6; hr=12; min=53; sec=5; ms=449;]

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Application No: 10576122 Version No: 3.0

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SEQUENCE LISTING

<110> MORGAN, Brian
 BURK, Mark
 LEVIN, Michael
 ZHU, Zoulin
 CHAPLIN, Jennifer
 KUSTEDJO, Karen
 HUANG, Zilin
 GREENBERG, William

<120> METHODS FOR MAKING SIMVASTATIN AND INTERMEDIATES

<130> 564462012800

<140> 10576122

<141> 2007-11-16

<150> PCT/US2004/034913

<151> 2004-10-20

<150> US 60/542,100

<151> 2004-02-04

<150> US 60/513,237

<151> 2003-10-21

<160> 6

<170> PatentIn version 3.1

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Gly Leu Glu Leu Pro Tyr Thr Thr Ile Thr Ser Ala Ala Val Ala Thr
 35 40 45

Glu Gly Pro Ile Pro Gln Pro Ala Ile Phe Gly Ser Thr Asp Pro Ile
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Val Ala Pro Glu Arg Cys Glu Val Arg Ala Val Thr Arg Pro Thr Lys
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Asp Ser Glu Ile Arg Ile Glu Leu Trp Leu Pro Leu Ser Gly Trp Asn
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Gly Lys Tyr Leu Gln Ile Gly Ser Gly Gly Trp Ala Gly Ser Ile Asn
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Arg Thr Gly Leu Ile Gly Pro Leu Gln Arg Gly Tyr Ala Val Ala Ala
 115 120 125

Thr Asp Asn Gly His Ile Ser Glu Gly Leu Val Pro Asp Ala Ser Trp
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Ala Ile Gly His Pro Gln Lys Leu Ile Asp Phe Gly Tyr Arg Ala Val
 145 150 155 160

His Glu Thr Ser Val Gln Ala Lys Ala Ile Leu Arg Ala Tyr Phe Gly
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Arg Gly Gln Asp Leu Ser Tyr Phe Ser Gly Cys Ser Asn Gly Gly Arg
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 195 200 205

Ile Ala Gly Ala Pro Ala Asn Asn Trp Ser Arg Leu Phe Thr Gly Phe
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Val Trp Asn Glu Arg Ala Leu Ala Asp Asp Pro Ile Pro Pro Ala Lys
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Gly Val Glu Asp Gly Leu Ile Glu Asn Pro Arg Ala Cys Ser Phe Asp
260 265 270

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Pro Arg Thr Gly Glu Arg Ile Phe Pro Gly Tyr Pro Met Gly Thr Glu
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Asp Val Ala Phe Gly Asp Ala Lys Ala Gly Pro Val Leu Asn Ala Thr
370 375 380

Asn Pro Asp Leu Arg Ser Phe Arg Ala Asn Gly Gly Lys Leu Ile Gln
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Tyr His Gly Trp Gly Asp Ala Ala Ile Thr Ala Phe Ser Ser Ile Asp
405 410 415

Tyr Tyr Glu Asn Val Arg Ala Phe Leu Asp Arg Phe Pro Asp Pro Arg
420 425 430

Ser Glu Asn Thr Asp Ile Asp Gly Phe Tyr Arg Leu Phe Leu Val Pro
435 440 445

Gly Met Gly His Cys Ser Gly Gly Ile Gly Pro Ser Ser Phe Gly Asn
450 455 460

Gly Phe Arg Ser Ala Arg Thr Asp Ala Glu His Asp Leu Leu Ser Ala

465 470 475 480

Leu Glu Ala Trp Val Glu Arg Asp Thr Ala Pro Glu Arg Leu Ile Gly
485 490 495

Thr Gly Thr Ala Val Gly Asp Pro Thr Ala Thr Leu Thr Arg Pro Leu
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<211> 1209

<212> DNA

<213> Unknown

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<223> Soil sample

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gtagtggaact tgtggggcgg catggcgcggt gccgacactc agacgccatg gacggcggag 180

acggtcagta ttgttttttc ctccaccaa ggcgcaacgg cactctgcgc ccatatgctg 240

gcgtcacgcg gccaaactgga tcttgatgca ccagtcgcca cctactggcc ggaatttgcc 300

caagccggca aagctcgcat cccggtgaaa atgctcttga accatcaagc tggctctcct 360

gccgtacgga caccgctgcc ccagggtgcc tacgctgact gggaaactgat ggtcaatacg 420

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ttccaagagg agatcgccag gccgttgggg ttagatttct ggattggctt accagcagag 600

caagaggcac gggtcgcgcc gatgatcgcg gcggagcctg atccgcaaag cctcttcttc 660

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Cys Val Thr Leu His Gly Glu Thr Val Val Asp Leu Trp Gly Gly Met
35 40 45

Ala Arg Ala Asp Thr Gln Thr Pro Trp Thr Ala Glu Thr Val Ser Ile
50 55 60

Val Phe Ser Ser Thr Lys Gly Ala Thr Ala Leu Cys Ala His Met Leu
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Ala Ser Arg Gly Gln Leu Asp Leu Asp Ala Pro Val Ala Thr Tyr Trp
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Pro Glu Phe Ala Gln Ala Gly Lys Ala Arg Ile Pro Val Lys Met Leu
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115

120

125

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130 135 140

Glu Pro Phe Trp Glu Pro Gly Thr Arg Asn Gly Tyr His Ala Leu Thr
145 150 155 160

Met Gly Trp Leu Val Gly Glu Val Val Arg Arg Val Ser Gly Lys Ser
165 170 175

Leu Gly Thr Phe Phe Gln Glu Glu Ile Ala Arg Pro Leu Gly Leu Asp
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Phe Trp Ile Gly Leu Pro Ala Glu Gln Glu Ala Arg Val Ala Pro Met
195 200 205

Ile Ala Ala Glu Pro Asp Pro Gln Ser Leu Phe Phe Gln Glu Val Ala
210 215 220

Lys Pro Gly Ala Leu Gln Ser Leu Val Leu Leu Asn Ser Gly Gly Tyr
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Met Gly Ala Gln Pro Glu Tyr Asp Ser Arg Ala Ala His Ala Ala Glu
245 250 255

Ile Gly Ala Ala Gly Gly Ile Thr Asn Ala Arg Gly Leu Ala Gly Met
260 265 270

Tyr Ala Pro Leu Ala Cys Gly Gly Lys Leu Lys Gly Val Glu Leu Val
275 280 285

Ser Pro Asp Met Leu Ala Arg Met Ser Arg Val Ala Ser Ala Thr Gly
290 295 300

Arg Asp Ala Val Leu Met Met Pro Thr Arg Phe Ala Leu Gly Phe Met
305 310 315 320

Lys Ser Met Asp Asn Arg Arg Glu Pro Ala Gly Val Gln Asp Ser Ala
325 330 335

Leu Phe Gly Glu Glu Ala Phe Gly His Val Gly Ala Gly Gly Ser Phe
340 345 350

Gly Phe Ala Asp Pro Lys Ala Gly Met Ser Phe Gly Tyr Thr Met Asn
 355 360 365

Arg Met Gly Leu Gly Ala Gly Leu Asn Pro Arg Gly Gln Ser Leu Val
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<212> DNA

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<221> SIGNAL

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35 40 45

Ala Gln Thr Val Thr Thr Gly Ser Leu Thr Pro Pro Gly Ser Thr Asn
50 55 60

Pro Ile Thr Asp Leu Pro Pro Phe Cys Arg Val Thr Gly Ala Ile Ala
65 70 75 80

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Thr	Ala	Ser	Thr	Asn	Thr	Gly	His	Glu	Ala	Ala	Pro	Gly	Met	Asn	Ala	130	135	140
Ala	Arg	Phe	Ala	Phe	Glu	Lys	Pro	Glu	Gln	Leu	Ile	Asp	Phe	Ala	Tyr	145	150	155
Arg	Ser	Gln	His	Glu	Thr	Ala	Leu	Lys	Ala	Lys	Ala	Leu	Val	Gln	Ala	165	170	175
Phe	Tyr	Gly	Lys	Pro	Pro	Glu	His	Ser	Tyr	Phe	Ile	Gly	Cys	Ser	Ser	180	185	190
Gly	Gly	Tyr	Gln	Gly	Leu	Met	Glu	Ala	Gln	Arg	Phe	Pro	Ala	Asp	Tyr	195	200	205
Asp	Gly	Ile	Val	Ala	Gly	Met	Pro	Ala	Asn	Asn	Trp	Thr	Arg	Leu	Met	210	215	220
Ala	Gly	Asp	Leu	Asp	Ala	Ile	Leu	Ala	Val	Ser	Val	Asp	Pro	Ala	Ser	225	230	235
His	Leu	Pro	Val	Ser	Ala	Leu	Gly	Leu	Leu	Tyr	Arg	Ser	Val	Leu	Ala	245	250	255
Ala	Cys	Asp	Gly	Ile	Asp	Gly	Val	Val	Asp	Gly	Val	Leu	Glu	Asp	Pro	260	265	270
Arg	Arg	Cys	Arg	Phe	Asp	Pro	Ala	Val	Leu	Met	Cys	Lys	Ala	Asp	Gln	275	280	285
Asn	Pro	Asp	Gly	Cys	Leu	Thr	Pro	Ala	Gln	Val	Glu	Ala	Ala	Arg	Arg	290	295	300

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Gly Leu Ala Pro Gly Ser Glu Pro Phe Trp Pro His Arg Asn Pro Ala
325 330 335

Asn Pro Phe Pro Ile Pro Ile Ala His Tyr Lys Trp Leu Val Phe Ala
340 345 350

Asp Pro Asn Trp As